IAP20 Rec'd PCT/PTO 15 JUN 2006

SU0423

SEQUENCE LISTING

```
<110> SUNTORY LIMITED
 <120> Arachidonic acid-containing plants and use of the plants
<130> SU0423
<140>
<141>
<150> JP 2003-419124
(151) 2003-12-17
<150> JP 2004-097089
(151) 2004-03-29
<160> 28
<170> Patentin Ver. 2.1
<210> 1
<211> 457
<212> PRT
<213> Mortierella alpina
Met Ala Ala Ala Pro Ser Val Arg Thr Phe Thr Arg Ala Glu lle Leu
1 5 10 15
Asn Ala Glu Ala Leu Asn Glu Gly Lys Lys Asp Ala Glu Ala Pro Phe
20 25 30
Leu Met Ile Ile Asp Asn Lys Val Tyr Asp Val Arg Glu Phe Val Pro
35 40 45
Asp His Pro Gly Gly Ser Val IIe Leu Thr His Val Gly Lys Asp Gly 50 60
Thr Asp Val Phe Asp Thr Phe His Pro Glu Ala Ala Trp Glu Thr Leu
65 70 75 80
Ala Asn Phe Tyr Val Gly Asp IIe Asp Glu Ser Asp Arg Ala IIe Lys
85 90 95
Asn Asp Asp Phe Ala Ala Glu Val Arg Lys Leu Arg Thr Leu Phe Gln 100 105 110
Ser Leu Gly Tyr Tyr Asp Ser Ser Lys Ala Tyr Tyr Ala Phe Lys Val
115 120 125
Ser Phe Asn Leu Cys IIe Trp Gly Leu Ser Thr Phe IIe Val Ala Lys
130 135 140
Trp Gly Gln Thr Ser Thr Leu Ala Asn Val Leu Ser Ala Ala Leu Leu
145 150 160
Gly Leu Phe Trp Gln Gln Cys Gly Trp Leu Ala His Asp Phe Leu His
165 170 175
His Gln Val Phe Gln Asp Arg Phe Trp Gly Asp Leu Phe Gly Ala Phe 180 185 190
Leu Gly Gly Val Cys Gln Gly Phe Ser Ser Ser Trp Trp Lys Asp Lys
195 200 205
His Asn Thr His His Ala Ala Pro Asn Val His Gly Glu Asp Pro Asp
210 215 220
lle Asp Thr His Pro Leu Leu Thr Trp Ser Glu His Ala Leu Glu Met 225 230 235 240
Phe Ser Asp Val Pro Asp Glu Glu Leu Thr Arg Met Trp Ser Arg Phe 245 250 255
```

 Met
 Val
 Leu
 Asn 260
 Gln
 Thr
 Trp
 Phe 286
 Phe Pro 11e
 SU0423 270
 Phe Ala

 Arg
 Leu
 Ser
 Trp
 Cys
 Leu
 Gln
 Ser
 Ile
 Met
 Phe
 Val
 Leu
 Pro
 Asn
 Gly

 Gln
 Ala
 His
 Lys
 Pro
 Ser
 Gly
 Ala
 Arg
 Val
 Pro
 Ile
 Pro
 Asn
 Gly

 Gln
 Leu
 Ser
 Leu
 Ala
 Met
 His
 Trp
 Tyr
 Tyr
 Tyr
 Leu
 Ala
 Thr
 Phe
 315
 Leu
 Asn
 Phe
 320

 Leu
 Phe
 Ile
 Lys
 Asp
 Pro
 Val
 Asn
 Met
 Ile
 Val
 Ile
 Phe
 320

 Leu
 Phe
 Ile
 Lys
 Asn
 Leu
 Leu
 Ala
 Ile
 Val
 Asn
 Phe
 Asn
 Phe

```
<210> 2
<211> 1371
<212> DNA
<213> Mortierella alpina
```

<400> 2 atggctgctg ctcccagtgt gaggacgttt actcgggccg agattttgaa tgccgaggcc 60 ctgaatgagg gcaagaagga tgccgaggca ccctttctga tgatcattga caacaaggtg 120 tacgatgtcc gcgagtttgt ccctgatcat cccggtggaa gtgtgattct cacgcacgtt 180 ggcaaggacg gcactgacgt ctttgacact ttccaccccg aggctgcttg ggagactctt 240 gccaactttt acgttggtga tattgatgag agcgatcgtg ccatcaagaa tgatgacttt 300 goggoogagg ttogcaagot gogcacottg ttocagtoco ttggotacta ogactogtoc 360 aaggcatact atgccttcaa ggtctcgttc aacctctgca tctgggggctt gtcgactttc 420 attgttgcca agtggggcca gacctcgacc ctogccaacg tgctctcggc tgcgctcttg 480 ggtctcttct ggcagcagtg cggatggttg gcgcacgact ttttgcacca ccaggtcttc 540 caggaccett totggggtga tottttoggo goottottgg gaggtgtotg coagggttto 600 togtoctoct ggtggaagga caagcacaac actoaccacg ctgctcccaa cgtccacggc 660 gaggatcccg acattgacac tcaccctctg ttgacctgga gtgagcatgc tctggagatg 720 tictoggatg ticotgacga ggagotgaco ogtatgtggt ogogottoat ggtoctoaac 780 cagacotggt totacitoco cattototog titgocogto tgtoctggtg cotocagico 840 attatgtttg ttctgcccaa cggtcaggcc cacaagccct ctggagcgcg tgtgcccatt 900 tegttggteg ageagetgte tetggetatg caetggaeet ggtacetege caecatgtte 960 etgtteatta aggateegt caacatgatt gtgtacettt tggtgtegea ggetgtttge 1020 ggcaacttgt tggcgattgt gttctcgctc aaccacaacg gcatgcctgt gatctccaag 1080 gaggaagcgg tcgatatgga cttcttcacc aagcagatca tcacgggtcg tgatgttcac 1140 cctggtctgt tigccaactg gitcacgggt ggattgaact accagaitga gcaccacttg 1200 ticccticga tgccccgcca caactitica aagatccagc ctgctgtcga gactitgtgc 1260 aaaaagtacg gtgtccgata ccataccact ggtatgatcg agggaactgc agaggtcttt 1320 agcogtttga acgaggtoto caaggoggoo tocaagatgg goaaggoaca g

```
<210> 3
<211> 318
<212> PRT
<213> Mortierella alpina
```

```
<400> 3
Met Glu Ser IIe Ala Gin Phe Leu Pro Ser Lys Met Pro Gin Asp Leu
1 10 15
Phe lle Asp Leu Ala Arg Ala lle Gly Val Gln Ala Ala Pro Tyr Val
20 25 30
Asp Pro Leu Glu Ala Ala Leu Val Ala Gin Ala Glu Lys Phe Phe Pro 35 40 45
Thr Val Val His His Thr Arg Gly Phe Leu Val Ala Val Glu Ser Pro
Leu Ala Arg Glu Leu Pro Leu Met Asn Pro Phe His Val Leu Leu IIe
65 70 75 80
Ala Leu Ala Tyr Leu Val Thr Val Phe Val Gly Met Gln lle Met Lys
85 90 95
Asn Phe Glu Arg Phe Glu Val Lys Thr Phe Ser Leu Phe His Asn Phe
100 105 110
Cys Leu Val Ser IIe Ser Ala Tyr Met Cys Gly Gly IIe Leu Tyr Glu
115 120 125
Ala Tyr Gln Ala Asn Tyr Gly Leu Phe Glu Asn Ala Ala Asp His Thr
130 140
Val Gln Gly Leu Pro Met Ala Lys Met IIe Trp Leu Phe Tyr Phe Ser
145 . 150 155 160
Lys lie Met Glu Phe Val Asp Thr Met lie Met Val Leu Lys Lys Asn
165 170 175
Asn Arg Gln Ile Ser Phe Leu His Val Tyr His His Ser Ser Ile Phe 180 185
Thr lie Trp Trp Leu Val Thr Phe Val Ala Pro Asn Gly Glu Ala Tyr 195 200 · 205
Phe Ser Ala Ala Leu Asn Ser Phe lle His Val lle Met Tyr Gly Tyr 210 215 220
Tyr Phe Leu Ser Ala Leu Gly Phe Lys Gln Val Ser Phe lle Lys Phe 225 230 235 240
Tyr lle Thr Arg Ser Gln Met Thr Gln Phe Cys Met Met Ser lle Gln 245 . 250 . 255
Ser Ser Trp Asp Met Tyr Ala Met Lys Val Leu Gly Arg Pro Gly Tyr 260 270
Pro Phe Phe IIe Thr Ala Leu Leu Trp Phe Tyr Met Trp Thr Met Leu 275 280 285
Gly Leu Phe Tyr Asn Phe Tyr Arg Lys Asn Ala Lys Leu Ala Lys Gln 290 295 300
Ala Lys IIe Asp Ala Ala Lys Glu Lys Ala Arg Lys Leu Gln 305 310 315
```

<210> 4 <211> 954

<212> DNA

<213> Mortierella alpina

<400> 4
atgagtcga ttgogcaatt cctcccctca aagatgccgc aagatctgtt tattgacctt 60
gcaagggcca tcggtgtcca ggccgcaccc tatgtcgacc ctctcgaggc agcgcttgtg 120
gccaggccg agaagttct ccccacggtc gttcatcaca cgcgcggctt tttggtcgcg 180
gtcgagtcac ccttggcccg tgagctgccc ttgatgaacc ccttccacgt gctgttgatc 240
gcgctcgctt acttggtcac ggtcttttgtg ggcatgcaga tcatgaagaa ctttgaacgg 300
ttcgaggtca agacgttct gctctccac aacttttgtc tggtctcgat cagtgcctac 360

SU0423

600

660

720

```
atgtgcggcg ggatcttgta cgaggcttac caggccaact atggactgtt tgagaacgcg
gccgatcata ccgtccaggg tcttcctatg gccaagatga tctggctctt ctacttctcc
aagatcatgg agtttgtcga caccatgatc atggtcctta agaagaacaa ccgccagatc
togttottgc acgtotacca coacagotoc atottoacca totggtggtt ggtoaccttt
gttgcaccca atggtgaagc ctacttctcg gctgcgttga actcgttcat ccacgtgatc
atgtacgget actacttect gteegeettg ggetteaage aggtgtegtt cateaagtte
tacatcacge gttcgcagat gacgcagtte tgcatgatgt cgatccagte etcetgggac 780 atgtatgcca tgaaggtget tggccgccc ggatacccet tettcatcae egcectgett 840 tggttctaca tgtggaccat geteggacte ttetacaact tetacagaaa gaacgccaag 900
ttggccaagc aggccaagat cgatgctgcc aaggagaagg caaggaagtt gcag
<210> 5
<211> 446
<212> PRT
<213> Mortierella alpina
 <400> 5
Met Gly Thr Asp Gln Gly Lys Thr Phe Thr Trp Gln Glu Leu Ala Ala
1 5 10
His Asn Thr Glu Asp Ser Leu Leu Leu Ala IIe Arg Gly Asn Val Tyr 25 30
Asp Val Thr Lys Phe Leu Ser Arg His Pro Gly Gly Thr Asp Thr Leu 35 40 45
Leu Leu Gly Ala Gly Arg Asp Val Thr Pro Val Phe Glu Met Tyr His
Glu Phe Gly Ala Ala Glu Ala Ile Met Lys Lys Tyr Tyr Val Gly Thr
65 70 75 80
Leu Val Ser Asn Glu Leu Pro IIe Phe Pro Glu Pro Thr Val Phe His
85 90 95
Lys Thr lie Lys Gly Arg Val Glu Ala Tyr Phe Lys Asp Arg Asn Met 100 105 110
Asp Ser Lys Asn Arg Pro Glu lle Trp Gly Arg Tyr Ala Leu lle Phe 115 120 125
Gly Ser Leu IIe Ala Ser Tyr Tyr Ala Gln Leu Phe Val Pro Phe Val
130 135 140
Val Glu Arg Thr Trp Leu Gln Val Val Phe Ala Ile Ile Met Gly Phe
145 150 160
Ala Cys Ala Gln Val Gly Leu Asn Pro Leu His Asp Ala Ser His Phe
165 170 175
Ser Val Thr His Asn Pro Thr Val Trp Lys IIe Leu Gly Ala Thr His
180 185 190
Asp Phe Phe Asn Gly Ala Ser Tyr Leu Val Trp Met Tyr Gln His Met 195 200 205
```

Phe Val Asn His IIe Asn Gln His Met Phe Val Pro Phe Leu Tyr Gly 245 250 255

Leu Gly His His Pro Tyr Thr Asn IIe Ala Gly Ala Asp Pro Asp Val 210 215 220

Ser Thr Ser Glu Pro Asp Val Arg Arg IIe Lys Pro Asn Gln Lys Trp 225 230 235 240

Leu Leu Ala Phe Lys Val Arg IIe Gin Asp IIe Asn IIe Leu Tyr Phe 260 265 270

Val Lys Thr Asn Asp Ala lle Arg Val Asn Pro lle Ser Thr Trp His 275 280 285

Thr Val Met Phe Trp Gly Gly Lys Ala Phe Phe Val Trp Tyr Arg Leu 290 295 300

```
lle Val Pro Met Gln Tyr Leu Pro Leu Ser Lys Val Leu Leu Leu Phe
305 310 320
Thr Val Ala Asp Met Val Ser Ser Tyr Trp Leu Ala Leu Thr Phe Gin 325 330 335
Ala Asn His Val Val Glu Val Gln Trp Pro Leu Pro Asp Glu Asn 340 345
Gly lle lle Gln Lys Asp Trp Ala Ala Met Gln Val Glu Thr Thr Gln 355 360 365
Asp Tyr Ala His Asp Ser His Leu Trp Thr Ser IIe Thr Gly Ser Leu
370 375 380
Asn Tyr Gln Ala Val His His Leu Phe Pro Asn Val Ser Gln His His 385 390 400
Tyr Pro Asp IIe Leu Ala IIe IIe Lys Asp Thr Cys Ser Glu Tyr Lys
405 410 415
Val Pro Tyr Leu Val Lys Asp Thr Phe Trp Gln Ala Phe Ala Ser His
420 425 430
Leu Glu His Leu Arg Val Leu Gly Leu Arg Pro Lys Glu Glu
435 440 445
<210> 6
<211> 1338
<212> DNA
<213> Mortierella alpina
<400> 6
atgggtacgg accaaggaaa aaccttcacc tggcaagaac tcgcggcgca taacaccgag 60 gacagcctcc ttttggctat ccgtggcaat gtatacgatg tcacaaagtt cttgagccgt 120
catcotggtg gaacggatac totottgctc ggagctggcc gagatgtcac tccggttttt 180
gagatgtace acgagttigg agetgeagag getateatga agaagtacta tgitgeaca 240 etggteteaa atgattge catetteeca gagecaacgg tgitecataa gaceateaag 300 ggeagagtig aggeatacti taaggacegg aacatggati ceaagaacag accagagate 360 tggggaegat atgeteeat ettiggatee tigategeet ettactaege geageteitt 420
gtaccgttcg tggtcgaacg tacatggctc caggtggtgt ttgctatcat catgggattt 480
gogtgogogo aagtoggatt gaaccetett cacgatgoet cocacttte agtgaccac 540 aaccecaceg titggaagat totoggagoo acgcacgact tittcaacgg agcatogtat 600
ctcgtgtgga tgtaccaaca tatgctcggc catcatccct ataccaacat tgctggagct 660
gatcocgatg tgtcgaccto tgagccogat gttcgtcgta tcaagcccaa ccaaaagtgg 720 ttcgtcaacc acatcaacca gcacatgttt gttcctttcc tgtacggact gctggcgttc 780 aaggtgcgca tccaggacat caacatcttg tactttgtca agaccaatga cgccattcgt 840 gtcaacccca tctcgacttg gcacaccgtc atgttctggg gcggaaaggc cttcttgtc 900 trgtacccat tgatcgttat tagacgtat atgacgatat caacactgt gcacaccgtc atgttctggg gcggaaaggc cttcttgtc 900
tggtaccgct tgatcgttcc tatgcagtat ctgcccctga gcaaggtgtt gctcttgttt 960
acceptogoag acategetot ticttacteg ctegogotega cottocaege gaaccaegtt 1020 gttgaggagg ttcagtggcc attgoctgat gagaateggaa toatcoaaaa ggattgggca 1080 gccategoagg togagactac tcaggattac gccacegatt cegoacctoteg gaccaegcatc 1140
acgggcaget tgaactacca agcogtteat catetgttee egaacgttte ceageateae 1200
taccotgata tootggotat catcaaggac acctgcagcg agtacaaggt gccatacctc 1260
gtcaaggata cottttggca agogtttgct toacatttgg agcacttgcg tgtgcttggt 1320
cttcgtccca aggaagag
<210> 7
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
         HinCprof
                                                                                                        26
agtcaagctt aattcaaaca aaaacg
```

<210> 8 <211> 26 <212> DNA

<213>	Artificial Sequence	
	Description of Artificial Sequence: Primer XbaCpror	
<400> cagtto	8 taga aaattottta atacgg	26
<210> <211> <212> <213>	26	
<220> <223>	Description of Artificial Sequence: Primer Sacmasf	
<400> agtcga	9 gctc cagcttccct gaaacc	26
<210> <211> <212> <213>	33	
<220> <223>	Description of Artificial Sequence: Primer Ecomasr	
<400> catcat	10 ctcg agggtggtga ccatggtgat cgc	33
<210> <211> <212> <213>	27	
	Description of Artificial Sequence: Primer det15-2-f1	
<400> atggtt	11 aaag acacaaagcc tttagcc	27
<210> <211> <212> <213>	22	
<220> <223>	Description of Artificial Sequence: Primer det15-2-r1	
<400> tcagtc	12 togt tgogagtgga gg	22
<210> <211> <212> <213>	33	
<220> <223>	Description of Artificial Sequence: Primer SOYF1-B	
<400> tggcct	13 ggga toottaaaga cacaaagoot tta	33
<210> <211> <212>	32	

(213) Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer SOYR1-X	
<pre><400> 14 gcacatctcg agggattgaa gtgagagcct tc</pre>	32
<210> 15 <211> 32 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer SOYF2-S	
<pre><400> 15 gtctgcgagc tcttaaagac acaaagcctt ta</pre>	32
<pre><210> 16 <211> 32 <212> DNA <213> Artificial Sequence</pre>	
<pre><220> <223> Description of Artificial Sequence: Primer SOUR2-X</pre>	
<400> 16 catcatctcg agggtggtga ccatggtgat gc	32
<210> 17 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer det6f3	
<400> 17 tggtggaagg acaagcacaa	20
<210> 18 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer det6r2	
<400> 18 acagaccagg gtgaacatca	20
<210> 19 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer det5f4	
<400> 19 ctttggatcc ttgatcgcct	20
<210> 20 <211> 20 <212> DNA <213> Artificial Sequence	
<220>	

	SU0423	
<223>	Description of Artificial Sequence: Primer det5r3	
<400> agaac	20 atgac ggtgtgccaa	20
<210> <211> <212> <213>	26	
<220> <223>	Description of Artificial Sequence: Primer XbaGLf	
<400> cagtt	21 ctaga goottotoac attooc	26
<210> <211> <212> <213>	26	
<220> <223>	Description of Artificial Sequence: Primer SacGLr	
<400> agtcg	22 agoto ttactgcaac ttoott	26
<210> <211> <212> <213>	20	
<220> <223>	Description of Artificial Sequence: Primer HPTf1	
<400> cctgc	23 gggta aatagotgog	20
<210> <211> <212> <213>	20	
<220> <223>	Description of Artificial Sequence: Primer HPTr1	
<400> cgtca	24 accaa gctctgatag	20
<210> <211> <212> <213>	20	
<220> <223>	Description of Artificial Sequence: Primer EGFP-f1	
<400> atggt	25 gagca agggcgagga	20
<210> <211> <212> <213>	21	
<220> <223>	Description of Artificial Sequence: Primer EGFP-R1	
<400>	26	

aatgaacatg togagcaggt a	SU0423
<210> 27 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	GLEf
<400> 27 gtgctcgctt atttggtcac	20
<210> 28 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	GLEr
<400> 28 cgacatcatg cagaactgtg	20